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Probabilistic Programming in neurolang: Bridging the Gap Between Cognitive Science and Statistical Modeling

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1 Introduction

Meta-analysis information often needs to be supplemented with neuroimaging or neuroanatomical data to build robust brain mapping models. To the best of our knowledge, a flexible tool for conducting this kind of analysis has yet to be proposed. We present neurolang: a convenient domain-specific language (DSL) for building rich brain mapping models. We demonstrate how this tool can be applied by combining meta-analysis data with atlas-based brain structures to construct anatomically-supported patterns of activation and reverse inference models.

Coordinate based meta-analysis (CBMA) tools, such as **Neurosynth** [Yarkoni et al., 2011], have become an integral part of brain mapping research. They proved very useful when combined with **functional magnetic resonance imaging (fMRI)** signals to derive activation patterns [e.g. Wager et al., 2013, Cole et al., 2012] or reveal meaningful cognitive processes through reverse inference [e.g. Smallwood and Schooler, 2015, Seghier, 2013, Chang et al., 2013, Andrews-Hanna et al., 2014]. Sometimes, within the same **fMRI** study, some **regions of interest (ROIs)** are built from meta-analysis forward inference maps while others are obtained from neuroanatomical probabilistic maps. This is because some **ROIs** are better supported by past literature while some others are better supported by neuroanatomy. There seems to be a space for a tool to define models that easily combine both modalities.

2 Methods

Adopting a language-oriented programming approach, we design a declarative and probabilistic **domain-specific language (DSL)** for formulating and testing sophisticated brain mapping hypotheses that benefit from mined meta-analysis and neuroanatomical data. A syntax close to natural language is translated to a set of Datalog rules that can also express probabilities. For example, "v is sensory if v is auditory or v is sensory if v is visual" translates to the datalog rule set $\{ \text{sensory}(v) :- \text{visual}(v), \text{sensory}(v) :- \text{auditory}(v) \}$ while "with probability p v is activated" translates to the ProbLog probabilistic fact $p :: \text{activated}(v)$, where p

could be a learnable parameter within a statistical model. Internally, intermediate representations based on established logic [Gallaire and Minker, 1978] and probabilistic programming [BRny et al., 2017] research are orchestrated to provide fast bayesian inference and maintain theoretical properties of programs.

3 Results

We present a real-world example originating from a study of the prefrontal cortex’s global connectivity [Cole et al., 2012] where ROIs are derived from both anatomical and meta-analysis data. The `neurolang` program in Figure 1 reconstructs these ROIs. A *sensory motor* ROI is derived by combining multiple cytoarchitecture probabilistic maps obtained using AFNI TT_N27_CA_EZ_PMaps. *Default mode network* and *cognitive control* ROIs are both derived from a forward inference meta-analysis model. Parameters of the model are defined as ProbLog probabilistic facts and estimated from Neurosynth data via a technique called “Learning From Interpretations” [Gutmann et al., 2011]. ROIs can then be obtained by formulating and solving queries on a graphical model translation of the inferred program. The language allows for a compact representation of the logical rules used to define these brain regions using multiple modalities. While in this example each region is based on a single modality, one could think of a region defined through a mixture of anatomical and meta-analytical probabilistic maps.

4 Conclusions

We develop a flexible and intuitive tool for formulating brain mapping hypotheses involving both neuroanatomical data and meta-analysis data. We present two examples involving both modalities to obtain accurate ROIs or terms most likely to correspond to a particular pattern of activation. This approach could help bridging the gap between computational neuroanatomy and rich statistical modeling. We acknowledge the support of ERC Neurolang.

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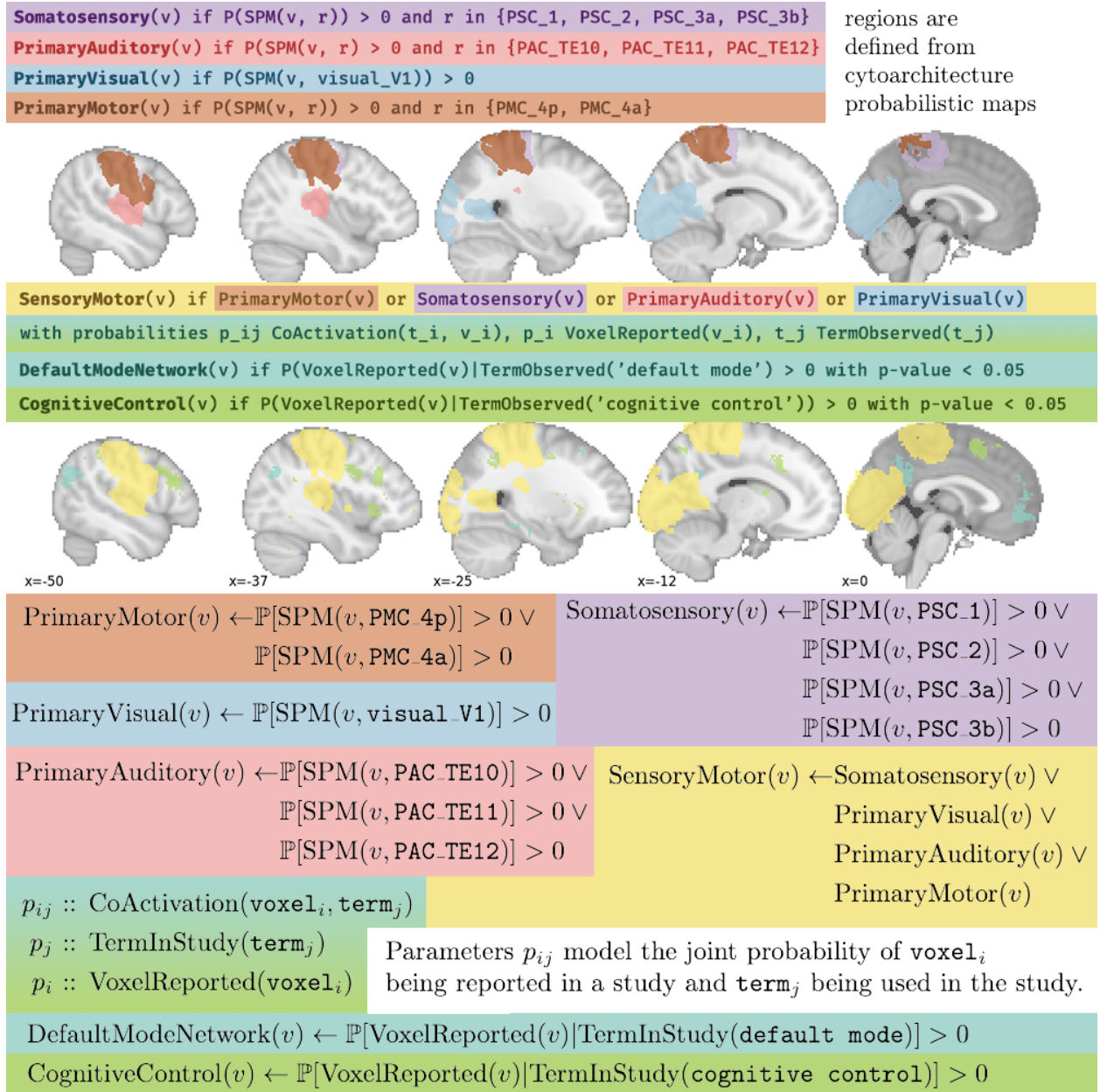


Figure 1: The program for constructing the sensory motor, default mode network and cognitive control ROIs is shown both in an intuitive DSL and in its logic programming intermediate representation. Colors are used to relate each rule of the program with its associated ROI in the brain plots. The construction of the *sensory motor* ROI is made iteratively by first constructing the somatosensory, primary auditory, primary visual and primary motor areas from the AFNI TT_N27_CA_EZ_PMaps probabilistic maps and then combining them. The *default mode network* and *cognitive control* ROIs are obtained by constructing a probabilistic model whose parameters are fitted on Neurosynth meta-analysis data. The obtained parameters are false discovery rate (FDR)-corrected for multiple comparisons using a whole-brain FDR threshold of 0.05, as in [Yarkoni et al., 2011].

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